## SEQUENCE LISTING

- (1) GENERAL INFORMATION:
  - (i) APPLICANTS: MUIR, TOM W.

MAYVILLE, PATRICIA NOVICK, RICHARD B. BEAVIS, RONALD JI, GUANGYONG

- (ii) TITLE OF INVENTION: NOVEL STAPHYLOCOCCUS PEPTIDES FOR BACTERIAL INTERFERENCE
- (iii) NUMBER OF SEQUENCES: 8
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: KLAUBER & JACKSON
  - (B) STREET: 411 HACKENSACK AVENUE
  - (C) CITY: HACKENSACK
  - (D) STATE: NEW JERSEY
  - (E) COUNTRY: USA
  - (F) ZIP: 07601
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Floppy disk
  - (B) COMPUTER: IBM PC compatible
  - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
  - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER: Not yet known
  - (B) FILING DATE: 24-JUN-1999
  - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
  - (A) APPLICATION NUMBER: 60/090,402
  - (B) FILING DATE: 24-JUN-1998
  - (C) CLASSIFICATION: Not assigned
- (viii) ATTORNEY/AGENT INFORMATION:
  - (A) NAME: JACKSON, DAVID A.
  - (B) REGISTRATION NUMBER: 26,742
  - (C) REFERENCE/DOCKET NUMBER: 600-1-231
  - (ix) TELECOMMUNICATION INFORMATION:
    - (A) TELEPHONE: 201-487-5800
    - (B) TELEFAX: 201-343-1684
- (2) INFORMATION FOR SEQ ID NO:1:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 9 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: not relevant
    - (D) TOPOLOGY: both
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Gly Val Asn Ala Xaa Ser Ser Leu Phe 1 5

- (2) INFORMATION FOR SEQ ID NO:2:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 9 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: not relevant
    - (D) TOPOLOGY: both
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
  - Gly Ala Asn Ala Xaa Ser Ser Leu Phe 1 5
- (2) INFORMATION FOR SEQ ID NO:3:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 9 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: not relevant
    - (D) TOPOLOGY: both
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
  - Gly Val Ala Ala Xaa Ser Ser Leu Phe 1 5
- (2) INFORMATION FOR SEQ ID NO:4:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 9 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: not relevant
    - (D) TOPOLOGY: both
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
  - Ala Val Asn Ala Xaa Ser Ser Leu Phe  $1 \hspace{1cm} 5$
- (2) INFORMATION FOR SEQ ID NO:5:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 9 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: not relevant
    - (D) TOPOLOGY: both
  - (ii) MOLECULE TYPE: peptide

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
- Gly Val Asn Ala Xaa Ala Ser Leu Phe
- (2) INFORMATION FOR SEQ ID NO:6:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 9 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: not relevant
    - (D) TOPOLOGY: both
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
  - Gly Val Asn Ala Xaa Ser Ala Leu Phe 5 1
- (2) INFORMATION FOR SEQ ID NO:7:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 9 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: not relevant
      (D) TOPOLOGY: both
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
  - Gly Val Asn Ala Xaa Ser Ser Ala Phe
- (2) INFORMATION FOR SEQ ID NO:8:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 5 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: not relevant
    - (D) TOPOLOGY: both
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

Xaa Ser Ser Leu Phe

## **SEQUENCE LISTING**

```
<110> Muir, Tom
   Mayville, Patricia
   Novick, Richard P.
   Beavis, Ronald
   Ji, Guangyong
<120> NOVEL STAPHYLOCOCCUS PEPTIDES FOR BACTERIAL
   INTERFERENCE
<130> 600-1-231N
<140> 09/339,511
<141> 1999-06-24
<150> 60/090,402
<151> 1998-06-24
<160>8
<170> Patentln Ver. 2.0
<210> 1
<211>9
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
   peptide
<220>
<221> VARIANT
<222> (5)
<223> Xaa represents any amino acid at this position.
<400> 1
Gly Val Asn Ala Xaa Ser Ser Leu Phe
<210>2
<211>9
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
   peptide
<220>
<221> VARIANT
<222> (5)
<223> Xaa represents any amino acid at this position.
```

```
<400> 2
 Gly Ala Asn Ala Xaa Ser Ser Leu Phe
            5
 <210>3
 <211>9
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Synthetic
    peptide
<220>
<221> VARIANT
<222> (5)
<223> Xaa represents any amino acid at this position.
<400> 3
Gly Val Ala Ala Xaa Ser Ser Leu Phe
<210>4
<211>9
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
   peptide
<220>
<221> VARIANT
<222> (5)
<223> Xaa represents any amino acid at this position.
<400> 4
Ala Val Asn Ala Xaa Ser Ser Leu Phe
 1
           5
<210>5
<211>9
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
   peptide
<220>
<221> VARIANT
<222> (5)
<223> Xaa represents any amino acid at this position.
```

```
<400>5
Gly Val Asn Ala Xaa Ala Ser Leu Phe
           5
<210>6
<211>9
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
   peptide
<220>
<221> VARIANT
<222> (5)
<223> Xaa represents any amino acid at this position.
<400>6
Gly Val Asn Ala Xaa Ser Ala Leu Phe
           5
<210>7
<211>9
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
   peptide
<220>
<221> VARIANT
<222> (5)
<223> Xaa represents any amino acid at this position.
<400> 7
Gly Val Asn Ala Xaa Ser Ser Ala Phe
           5
<210>8
<211>5
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
   peptide
<220>
<221> VARIANT
<222> (1)
```

<223> Xaa represents any amino acid at this position.

<400> 8 Xaa Ser Ser Leu Phe 1 5